SEQUENCE LISTING

<110> Tang, Y. Tom
Guegler, Karl J.
Corley, Neil C.
Gorgone, Gina A.
Yue, Henry

<120> CALCIUM BINDING PROTEIN

<130> PF-0635-2 DIV

<140> To Be Assigned
<141> Herewith

<160> 5

<170> PERL Program

<210> 1
<211> 337
<212> PRT
<213> Homo sapiens

<220> -
<223> 3734805

<400> 1

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Glu Ile Val Lys Ile Leu Lys Asp Asn Leu Ala Ile Leu Glu Lys
             20                  25                  30
Gln Asp Lys Lys Thr Asp Lys Ala Ser Glu Glu Val Ser Lys Ser
         35                  40                  45
Leu Gln Ala Met Lys Glu Ile Leu Cys Gly Thr Asn Glu Lys Glu
     50                  55                  60
Pro Pro Thr Glu Ala Val Ala Gln Leu Ala Gln Glu Leu Tyr Ser
 65                  70                  75
Ser Gly Leu Leu Val Thr Leu Ile Ala Asp Leu Gln Leu Ile Asp
             80                  85                  90
Phe Glu Gly Lys Lys Asp Val Thr Gln Ile Phe Asn Asn Ile Leu
         95                 100                 105
Arg Arg Gln Ile Gly Thr Arg Ser Pro Thr Val Glu Tyr Ile Ser
    110                 115                 120
Ala His Pro His Ile Leu Phe Met Leu Leu Lys Gly Tyr Glu Ala
    125                 130                 135
Pro Gln Ile Ala Leu Arg Cys Gly Ile Met Leu Arg Glu Cys Ile
    140                 145                 150
Arg His Glu Pro Leu Ala Lys Ile Ile Leu Phe Ser Asn Gln Phe
    155                 160                 165
Arg Asp Phe Phe Lys Tyr Val Glu Leu Ser Thr Phe Asp Ile Ala
    170                 175                 180
Ser Asp Ala Phe Ala Thr Phe Lys Asp Leu Leu Thr Arg His Lys
    185                 190                 195
Val Leu Val Ala Asp Phe Leu Glu Gln Asn Tyr Asp Thr Ile Phe
    200                 205                 210
Glu Asp Tyr Glu Lys Leu Leu Gln Ser Glu Asn Tyr Val Thr Lys
    215                 220                 225
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PF-0635-2 DIV

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Arg Gln Ser Leu Lys Leu Leu Gly Glu Leu Ile Leu Asp Arg His
                230                 235                 240
Asn Phe Ala Ile Met Thr Lys Tyr Ile Ser Lys Pro Glu Asn Leu
                245                 250                 255
Lys Leu Met Met Asn Leu Leu Arg Asp Lys Ser Pro Asn Ile Gln
                260                 265                 270
Phe Glu Ala Phe His Val Phe Lys Val Phe Val Ala Ser Pro His
                275                 280                 285
Lys Thr Gln Pro Ile Val Glu Ile Leu Leu Lys Asn Gln Pro Lys
                290                 295                 300
Leu Ile Glu Phe Leu Ser Ser Phe Gln Lys Glu Arg Thr Asp Asp
                305                 310                 315
Glu Gln Phe Ala Asp Glu Lys Asn Tyr Leu Ile Lys Gln Ile Arg
                320                 325                 330
Asp Leu Lys Lys Thr Ala Pro
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<210> 2
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<212> DNA
<213> Homo sapiens

<220> -
<223> 3734805

<400> 2

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atcctgaaag acaatttggc cattttggaa aagcaagaca aaaagacaga caaggcttca 240
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gaaccccga cagaagcagt ggctcagcta gcacaagaac tctacagcag tggcctgctg 360
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atatttaaca acatcttgag aagacagata ggcactcgga gtcctactgt ggagtatatt 480
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<212> PRT
<213> Mus sp.

<220> -

PF-0635-2 DIV

<223> g262934

<400> 3

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             20                  25                  30
Ile Ser Asp Lys Lys Ala Glu Lys Ala Thr Glu Glu Val Ser Lys
             35                  40                  45
Asn Leu Val Ala Met Lys Glu Ile Leu Tyr Gly Thr Asn Glu Lys
             50                  55                  60
Glu Pro Gln Thr Glu Ala Val Ala Gln Leu Ala Gln Glu Leu Tyr
             65                  70                  75
Asn Ser Gly Leu Leu Gly Thr Leu Val Ala Asp Leu Gln Leu Ile
             80                  85                  90
Asp Phe Glu Gly Lys Lys Asp Val Ala Gln Ile Phe Asn Asn Ile
             95                 100                 105
Leu Arg Arg Gln Ile Gly Thr Arg Thr Pro Thr Val Glu Tyr Ile
            110                 115                 120
Cys Thr Gln Gln Asn Ile Leu Phe Met Leu Leu Lys Gly Tyr Glu
            125                 130                 135
Ser Pro Glu Ile Ala Leu Asn Cys Gly Ile Met Leu Arg Glu Cys
            140                 145                 150
Ile Arg His Glu Pro Leu Ala Lys Ile Ile Leu Trp Ser Glu Gln
            155                 160                 165
Phe Tyr Asp Phe Phe Arg Tyr Val Glu Met Ser Thr Phe Asp Ile
            170                 175                 180
Ala Ser Asp Ala Phe Ala Thr Phe Lys Asp Leu Leu Thr Arg His
            185                 190                 195
Lys Leu Leu Ser Ala Glu Phe Leu Glu Gln His Tyr Asp Arg Phe
            200                 205                 210
Phe Ser Glu Tyr Glu Lys Leu Leu His Ser Glu Asn Tyr Val Thr
            215                 220                 225
Lys Arg Gln Ser Leu Lys Leu Leu Gly Glu Leu Leu Leu Asp Arg
            230                 235                 240
His Asn Phe Thr Ile Met Thr Lys Tyr Ile Ser Lys Pro Glu Asn
            245                 250                 255
Leu Lys Leu Met Met Asn Leu Leu Arg Asp Lys Ser Arg Asn Ile
            260                 265                 270
Gln Phe Glu Ala Phe His Val Phe Lys Val Phe Val Ala Asn Pro
            275                 280                 285
Asn Lys Thr Gln Pro Ile Leu Asp Ile Leu Leu Lys Asn Gln Thr
            290                 295                 300
Lys Leu Ile Glu Phe Leu Ser Lys Phe Gln Asn Asp Arg Thr Glu
            305                 310                 315
Asp Glu Gln Phe Asn Asp Glu Lys Thr Tyr Leu Val Lys Gln Ile
            320                 325                 330
Arg Asn Leu Lys Arg Ala Ala Gln Gln Glu Ala
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<212> PRT
<213> Drosophila melanogaster

<220> -
<223> g1794137

PF-0635-2 DIV

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Lys Val Glu Lys Ala Gln Glu Asp Val Ser Lys Asn Leu Val Ser
35 40 45
Ile Lys Asn Met Leu His Gly Ser Ser Asp Ala Glu Pro Pro Ala
50 55 60
Asp Tyr Val Val Ala Gln Leu Ser Gln Glu Leu Tyr Asn Ser Asn
65 70 75
Leu Leu Leu Leu Leu Ile Gln Asn Leu His Arg Ile Asp Phe Glu
80 85 90
Gly Lys Lys His Val Ala Leu Ile Phe Asn Asn Leu Leu Arg Arg
95 100 105
Gln Ile Gly Thr Arg Ser Pro Thr Val Glu Tyr Ile Cys Thr Lys
110 115 120
Pro Glu Ile Leu Phe Thr Leu Met Ala Gly Tyr Glu Asp Ala His
125 130 135
Pro Glu Ile Ala Leu Asn Ser Gly Thr Met Leu Arg Glu Cys Ala
140 145 150
Arg Tyr Glu Ala Leu Ala Lys Ile Met Leu His Ser Asp Glu Phe
155 160 165
Phe Lys Phe Phe Arg Tyr Val Glu Val Ser Thr Phe Asp Ile Ala
170 175 180
Ser Asp Ala Phe Ser Thr Phe Lys Glu Leu Leu Thr Arg His Lys
185 190 195
Leu Leu Cys Ala Glu Phe Leu Asp Ala Asn Tyr Asp Lys Phe Phe
200 205 210
Ser Gln His Tyr Gln Arg Leu Leu Asn Ser Glu Asn Tyr Val Thr
215 220 225
Arg Arg Gln Ser Leu Lys Leu Leu Gly Glu Leu Leu Leu Asp Arg
230 235 240
His Asn Phe Thr Val Met Thr Arg Tyr Ile Ser Glu Pro Glu Asn
245 250 255
Leu Lys Leu Met Met Asn Met Leu Lys Glu Lys Ser Arg Asn Ile
260 265 270
Gln Phe Glu Ala Phe His Val Phe Lys Val Phe Val Ala Asn Pro
275 280 285
Asn Lys Pro Lys Pro Ile Leu Asp Ile Leu Leu Arg Asn Gln Thr
290 295 300
Lys Leu Val Asp Phe Leu Thr Asn Phe His Thr Asp Arg Ser Glu
305 310 315
Asp Glu Gln Phe Asn Asp Glu Lys Ala Tyr Leu Ile Lys Gln Ile
320 325 330
Lys Glu Leu Lys Pro Leu Pro Glu Ala
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<210> 5
<211> 377
<212> PRT
<213> Caenorhabditis elegans

<220> -
<223> g1255838

<400> 5

PF-0635-2 DIV

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Val Lys Thr Leu Arg Glu Val Leu Thr Ile Leu Asp Lys Leu Pro
             20                  25                  30
Pro Pro Lys Leu Asp Lys Asp Gly Asn Ile Gln Ser Asp Lys Lys
         35                  40                  45
Tyr Asp Lys Ala Leu Asp Glu Val Ser Lys Asn Val Ala Met Ile
     50                  55                  60
Lys Ser Phe Ile Tyr Gly Asn Asp Ser Ala Glu Pro Ser Ser Glu
 65                  70                  75
His Val Val Gln Val Ala Gln Leu Ala Gln Glu Val Tyr Asn Ala
             80                  85                  90
Asn Ile Leu Pro Met Leu Ile Lys Met Leu Pro Lys Phe Glu Phe
         95                 100                 105
Glu Cys Lys Lys Asp Val Gly Gln Ile Phe Asn Asn Leu Leu Arg
    110                 115                 120
Arg Gln Ile Gly Thr Arg Ser Pro Thr Val Glu Tyr Leu Gly Ala
    125                 130                 135
Arg Pro Glu Ile Leu Ile Gln Leu Val Gln Gly Tyr Ser Val Pro
    140                 145                 150
Asp Ile Ala Leu Thr Cys Gly Leu Met Leu Arg Glu Ser Ile Arg
    155                 160                 165
His Asp His Leu Ala Lys Ile Ile Leu Tyr Ser Asp Val Phe Tyr
    170                 175                 180
Thr Phe Phe Leu Tyr Val Gln Ser Glu Val Phe Asp Ile Ser Ser
    185                 190                 195
Asp Ala Phe Ser Thr Phe Lys Glu Leu Thr Thr Arg His Lys Ala
    200                 205                 210
Ile Ile Ala Glu Phe Leu Asp Ser Asn Tyr Asp Thr Phe Phe Ala
    215                 220                 225
Gln Tyr Gln Asn Leu Leu Asn Ser Lys Asn Tyr Val Thr Arg Arg
    230                 235                 240
Gln Ser Leu Lys Leu Leu Gly Glu Leu Leu Leu Asp Arg His Asn
    245                 250                 255
Phe Asn Thr Met Thr Lys Tyr Ile Ser Asn Pro Asp Asn Leu Arg
    260                 265                 270
Leu Met Met Glu Leu Leu Arg Asp Lys Ser Arg Asn Ile Gln Tyr
    275                 280                 285
Glu Ala Phe His Val Phe Lys Val Phe Val Ala Asn Pro Asn Lys
    290                 295                 300
Pro Lys Pro Ile Ser Asp Ile Leu Asn Arg Asn Arg Glu Lys Leu
    305                 310                 315
Val Glu Phe Leu Ser Glu Phe His Asn Asp Arg Thr Asp Asp Glu
    320                 325                 330
Gln Phe Asn Asp Glu Lys Ala Tyr Leu Ile Lys Gln Ile Gln Glu
    335                 340                 345
Met Lys Ser Ser Pro Lys Glu Ala Lys Lys Pro Lys Ser Lys Glu
    350                 355                 360
Asp Glu Asn Gln Glu Pro Ala Gly Pro Ser Glu Gly Pro Ser Thr
    365                 370                 375
Ser Gln
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